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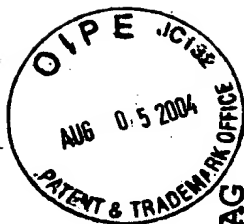
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181 C C C G G T C G A C G C C T G G G T C T G C C G G A A G A G C G A T G A G A G G T G T C T G A A G G T G G C T A T
241 T C A C T G A C G C A T G G G G T T G G A C T T G A A G G A T G C C A A G A G A T G C T G C C C C C A C C C C C T T A

1 M G P E A L S S L L L L L L
301 G G C C C G A G G A T C A G G A G C T A T G G G A C C A G A G C C C T G T C A T C T T T A C T G C T G C T G C T C T

15 V A S G D A D M K G H F D P A K C R Y A
361 T G G T G G C A A G T G G A G A T G C T G A C A T G A A G G G A C A T T T T G A T C C T G C C A A G T G C C G C T A T G

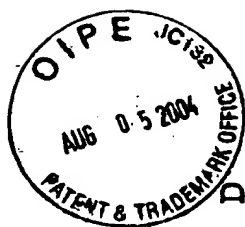
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421 C C C T G G G C A T G C A G G A C C G G A C C A T C C C A G A C A G T G A C A T C T C T G C T T C C A G C T C C T G G T

55 D S T A A R H S R L E S S D G D G A W C
481 C A G A T C C A C T G C C G C C C C A C A G C A G G T T G G A G A G C A G T G A C G G G G A T G G G C C T G G T

75 P A G S V F P K E E Y L Q V D L Q R L
541 G C C C C G A G G T C G G T G T T T C C C A A G A G A G G A G T A C T T G C A G G T G G A T C T A C A C G A C

95 H L V A L V G T Q G R H A G G L G K E F
601 T C C A C C T G G T G G C T C T G G T G G C A C C C A G G A C G G C A T G C C G G G G C C T G G G C A A G G A G T

FIG.1A



115 S R S Y R L R Y S R D G R R W M G W K D
661 TCTCCGGAGCTACCGCTGCGTTACTCCCGGATGTCGCCGCTGGATGGCTGGAAGG
135 R W G Q E V I S G N E D P E G V V L K D
721 ACCGTGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCCTGAGGGAGTGGTGTGAAGG
155 L G P P M V A R L V R F Y P R A D R V M
781 ACCTTGGCCCCCATGTTGCCCGACTGGTTCGCTTCTACCCCGGCTGACCGGTCA
175 S V C L R V E L Y G C L W R D G L L S Y
841 TGAGTGTCTGTCGGGTAGAGCTCTATGGCTGCCCTCTGGAGGATGGACTCCTGTCTT
195 T A P V G Q T M Y L S E A V Y L N D S T
901 ACACCGCCCTGTGGGCAGACAATGTATTATCTGAGGCCGTGTACCTCAACGACTCCA
215 Y D G H T V G G L Q Y G G L G Q L A D G
961 CCTATGACGGACATACCGTGGCGGACTGCAGTATGGGGTCTGGGCCAGCTGGCAGATG
235 V V G L D D F R K S Q E L R V W P G Y D
1021 GTGTGGTGGGCTGGATGACTTTAGGAAGAGTCAGGAGCTCGGGTCTGGCCAGGCTATG
255 Y V G W S N H S F S S G Y V E M E F E F
1081 ACTATGTGGATGGAGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGT

FIG.1B



275 D R L R A F Q A M Q V H C N N M H T L G
1141 TTGACCGGCTGAGGCCCTTCCAGGCTATGCAGGTCCACTGTAAACAATGCACACGCTGG
295 A R L P G G V E C R F R R G P A M A W E
1201 GAGCCCGTCTGCCCTGGCGGGTGAATGCTCGCTTCCGGCGTGCCCTGCCATGGCCTGGG
315 G E P M R H N L G G N L G D P R A R A V
1261 AGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTGGGGACCCAGAGCCCGGGCTG
335 S V P L G G R V A R F L Q C R F L F A G
1321 TCTCAGTGCCCGCTTGGCGCGTGCTGGCTTCTGCAGTGCCGCTTCTCTTTGCGG
355 P W L L F S E I S F I S D V V N N S S P
1381 GGCCCTGGTTACTCTTCAGCGAAATCTCCTTCACTCTGATGTGTGAACAATTCCTCTC
375 A L G G T F P P A P W P P G P P P T N
1441 CGGCACTGGGAGGCACCTTCCCGCAGCCCGCTGGTGCCGCTGGCCACCTCCACCA
395 F S S L E L E P R G Q Q P V A K A E G S
1501 ACTTCAGCAGCTTGGAGCTGGAGCCAGAGCCAGCAGCCCGTGCCCAAGGCCGAGGGA
415 P T A I L I G C L V A I I L L L L I I
1561 GCCGACCGCCATCCTCATCGGCTGCCCTGGTGGCCATCATCTGCTCTGCTGCTCATCA

FIG.1C



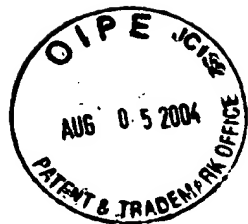
435 A L M L W R L H W R R L L S K A E R R V
1621 TTGCCCTCATGCTCTGGGGCTGCACTGGCGAGGCTCCTCAGCAAGGCTGAACGGAGGG
455 L E E E L T V H L S V P G D T I L I N N
1681 TGTGGAAGAGGAGCTGACGGTTACCTCTCTGTCCCTGGGACACTATCCTCATCAACA
475 R P G P R E P P P Y Q E P R P R G N P P
1741 ACCGCCAGGTCCTAGAGAGCCACCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGC
495 H S A P C V P N G S A L L L S N P A Y R
1801 CCCACTCCGCTCCCTGTGTCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACC
515 L L L A T Y A R P P R G P G P P T P A W
1861 GCCTCCTTCTGGCCACTTACGCCCTCCCTCGAGGCCCGGGCCCCCACACCCGCCCT
535 A K P T N T Q A Y S G D Y M E P E K P G
1921 GGGCCAAACCCACCAACACCCAGGCCCTACAGTGGGACTATATGGAGCCTGAGAAGCCAG
555 A P L L P P P P Q N S V P H Y A E A D I
1981 GCGCCCCGCTTCTGCCCCCACCCTCCCGAAGACAGCGTCCCCCATTTATGCCGAGGCTGACA
575 V T L Q G V T G G N T Y A V P A L P P G
2041 TTGTTACCCCTGCAGGGGCTACCGGGGCAACACCTATGCTGTGCTGCTGCACTGCCCCCCAG

FIG.1D



595 A V G D G P P R V D F P R S R L R F K E
2101 GGCAGTCGGGATGGCCCCCAGAGTGGATTTCCTCGATCTCGACTCCGCTCAAGG
615 K L G E G Q F G E V H L C E V D S P Q D
2161 AGAAGCTTGGCGAGGCCAGTTTGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAG
635 L V S L D F P L N V R K G H P L L V A V
2221 ATCTGGTCAGTCTTGATTTCCTCCCTTAATGTGCGTAAGGGACACCCCTTTGCTGGTAGCTG
655 K I L R P D A T K N A S F S L F S R N D
2281 TCAAGATCTTACGGCCAGATGCCACCAAGAAATGCCAGCTTCTCCTTGTCTCCAGGAATG
675 F L K E V K I M S R L K D P N I I R L L
2341 ATTTCTGAAAGAGGTGAAGATCATGTGCGAGGCTCAAGGACCCCAACATCATTCGGCTGC
695 G V C V Q D D P L C M I T D Y M E N G D
2401 TGGCGTGTGTGTCAGGACGACCCCTCTGTCATGATTACTGACTACATGGAGAACGGCG
715 L N Q F L S A H Q L E D K A A E G A P G
2461 ACCTCAACCAGTTCTCAGTGGCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCCTG
735 D G Q A A Q G P T I S Y P M L L H V A A
2521 GGGACGGCAGGCTGCGCAGGGGCCACCATCAGCTACCCAATGCTGCTGTCATGTGGCAG

FIG.1E



755 Q I A S G M R Y L A T L N F V H R D L A
2581 CCCAGATCGCCTCCGGCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGG
775 T R N C L V G E N F T I K I A D F G M S
2641 CCACGCGAACTGCCTAGTTGGGAAATTTACCATCAAAATCGCAGACTTTGGCATGA
795 R N L Y A G D Y Y R V Q G R A V L P I R
2701 GCCGGAACCTCTATGCTGGGACTATTACCGTGTGCGAGGCCGGCAGTGCTGCCCCATCC
815 W M A W E C I L M G K F T T A S D V W A
2761 GCTGGATGCCCTGGAGTGTCATCTCATGGGAAAGTTACGACTGCGAGTGACGTGTGGG
835 F G V T L W E V L M L C R A Q P F G Q L
2821 CCTTTGGTGTGACCCCTGTGGAGGTGCTGATGCTCTGTAGGGCCAGCCCTTTGGGCAGC
855 T D E Q V I E N A G E F F R D Q G R Q V
2881 TCACCGACGAGCAGGTTCATCGAGAACCGGGGGAGTTCTTCCGGGACCAGGGCCGGCAGG
875 Y L S R P P A C P Q G L Y E L M L R C W
2941 TGTACCTGTCCCGCCCTGCTGCCCCGAGGGCCATATGAGCTGATGCTTCGGTGCT
895 S R E S E Q R P P F S Q L H R F L A E D
3001 GGAGCCGGAGTCTGAGCAGCAGCACCCCTTTTCCAGCTGCATCGGTTCTCTGGCAGAGG

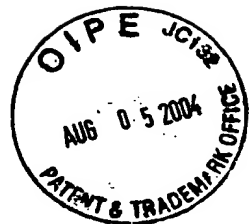
FIG.1F



915 A L N T V
3061 ATGCACTCAACACGGTGTGAATCACACATCCAGCTGCCCCCTCCCTCAGGGAGTGATCCAG

3121 GGAAGCCAGTGACACTAAACAAGAGGACACAATGGCACCTCTGCCCCCTCCCCCTCCCGA
3181 CAGCCCATCACCTCTAATAGAGGCAGTGAGACTGCAGGTGGCTGGGCCACCCAGGGAG
3241 CTGATGCCCTTCTCCCTTCCCTGGACACACTCTCATGTCTCCCTTCTCTTCTTCTTCC
3301 TAGAAGCCCCCTGTGCGCCCAACCCAGCTGGTCTGTGGATGGGATCCTCTCCACCCCTCTCT
3361 AGCCATCCCTTGGGAAGGTGGGAGAAATATAGGATAGACACTGGACATGGCCCATTTG
3421 GAGCACCTGGGCCCCACTGGACACACACTGATTCCTGGAGAGGTGGCTGCGCCCAAGCTTC
3481 TCTCTCCCTGTACACACACTGGACCCCACTGGCTGAGAACTCTGGGGGTGAGGAGGACAAGA
3541 AGGAGAGGAATAATGTTTCTTGTGCTGCTGCTCTGTACTTGTCTCAGCTTGGGCTTCTTC
3601 CTCCTCCATCACCTGAACACACTGGACCTGGGGGTAGCCCCGCCCAAGCTCAGTCAACCC
3661 CCACCTCCCACTTGCAGCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAG
3721 TAAATATTGGGATTGGGGGAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATC
3781 TCTAGTGTAGCTGCCACATTTGATTTTCTATAATCATCTTGGGGTTGTACATTTTGGGG
3841 GGAGAGACACAGATTTTACACTAATAATATGGACCTAGCTTGAGGCAATTTTAATCCCCCT
3901 GCACTAGGCAGGTAATAATAAAGGTGAGTTTTCACAAATAAATAAATAAATAAATAAATAA
3961 TC

FIG.1G



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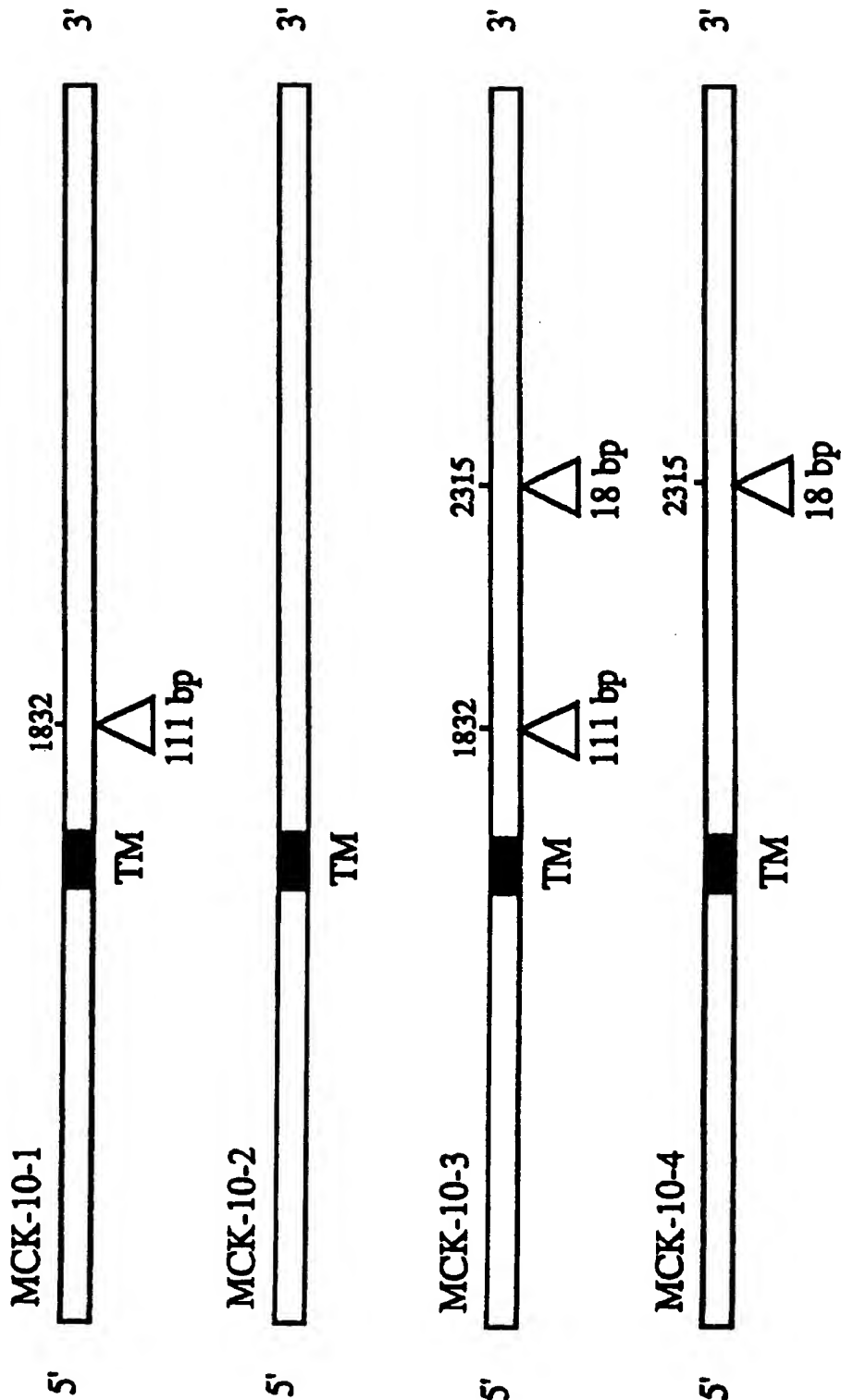


FIG.2

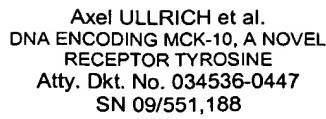


FIG.3A



542 gcccaatgatggaaggctggactcagaagaagggatggagcctggcgcclgagatccca 601
cggtttataccctccgacctgagctctctccctaccctcggaccacgggactctaaaggt
b A K Y G R L D S E E G D G A W C P E I P -
602 gtggaacctgatgacctgaaggagttctgcagattgacctgcacacctccattttatc 661
caccttggactactggacttccctcaaaagcgtctaaactgaacgtgtggggaggtaaaatag
b V E P D D L K E F L Q I D L H T L H F I -
662 actctgggtggggaccagggggcgccgagcaggaggctcatggcatcgagtttgcctccatg 721
tgagaccaccttgggtccctcgggctcgtctccagctaccgtagctcaaacgggggtac
b T L V G T Q G R R A G G H G I E F A P M -
722 tacaagatcaattacagctcgggatggcactcgtctggatctcttggcggaaacctcatggg 781
atgttctagttaatgtcagccctaccgtagcggacctagagaacctgccttggcagctacc
b Y K I N Y S R D G T R W I S W R N R H G -
762 aaacagggtgctggatggaaatagtaacctatgacatttcttaaggaccttggagccg 841
tttgcctcagacctaccctttatcatlggggatctgtaaaaggatttcttgaacctcggc
b K Q V L D G N S N P Y D I F L K D L E P -
842 cccattgtagccagatttgcctgggttcatlccagctaccgaccttccatgaatgttgt 901
gggtaacatcgggtctaaacaggccaagttaaggctcagtggtgggtgaggtacttccacaca
b P I V A R F V R F I P V T D H S M N V C -
902 atgagagtgaggctttacggCTGTGTCTGCCTAGATGCCTTGGTGTCTTACAATGCTCCA 961
tactctcaccctcgaootgccGACACAGACCGATCTACCGAACCACAGAATGTTACGAGGT
b M R V E L Y G C V W L D G L V S Y N A P -

FIG.3B



962 GCTGGGCAGCAGTTTGTACTCCCTGGAGGTTCCATCATTATCTGAATGATTCTGTCTAT 1021
CGACCCGTCGTCAAACATGACGGACCTCCAAGGTAGTAAATAGACTTACTAAGACAGATA

b A G Q Q F V L P G G S I I Y L N D S V Y -

1022 GATGGAGCTGTTGGATACAGCATGACAGAAGGGCTAGGCCAATTGACCGATGGTGTGTCT 1081
CTACCTCGACAACCTATGTCGTACTGTCTTCCCGATCCGGTAACTGGCTACACACAGA

b D G A V G Y S M T E G L G Q L T D G V S -

1082 GGCCTGGACGATTTACCCAGACCCATGAATACCACGTGTGGCCGGCTATGACTATGTG 1141
CCGGACCTGCTAAAGTGGTCTGGGTACTTATGGTGCACACGGGCGGATACTGATACAC

b G L D D F T Q T H E Y H V W P G Y D Y V -

1142 GGCTGGCGGAACGAGAGTGCCACCAATGGCTACATTGAGATCATGTTTGAATTTGACCGC 1201
CCGACCGCCTTGCTCTCACGGTGGTTACCGATGTAACCTAGTACAACTTAACTGGCG

b G W R N E S A T N G Y I E I M F E F D R -

1202 ATCAGGAATTTCACTACCATGAAGGTCCACTGCAACAACATGTTTGCTAAAGGTGTGAAG 1261
TAGTCCTTAAAGTGATGGTACTTCCAGGTGACGTTGTTGTACAAACGATTTCCACACTTC

b I R N F T T M K V H C N N M F A K G V K -

1262 ATCTTTAAGGAGGTACAGTGCTACTTCCGCTCTGAAGCCAGTGAGTGGGTACCTAATGCC 1321
TAGAAATTCCTCCATGTCACGATGAAGGCGAGACTTCGGTCACTCACCCATCGATTACGG

b I F K E V Q C Y F R S E A S E W V P N A -

1322 ATTCCTTcccccttgtcctggatgacgtcaoccccgltgctcggttltgtcacggltgcct 1381
TAAAGGAagggggaocaggacctoctgcogltggggtcacgagcccaocogltgccacgga

b I S F P L V L D D V N P S A R F V T V P -

FIG.3C

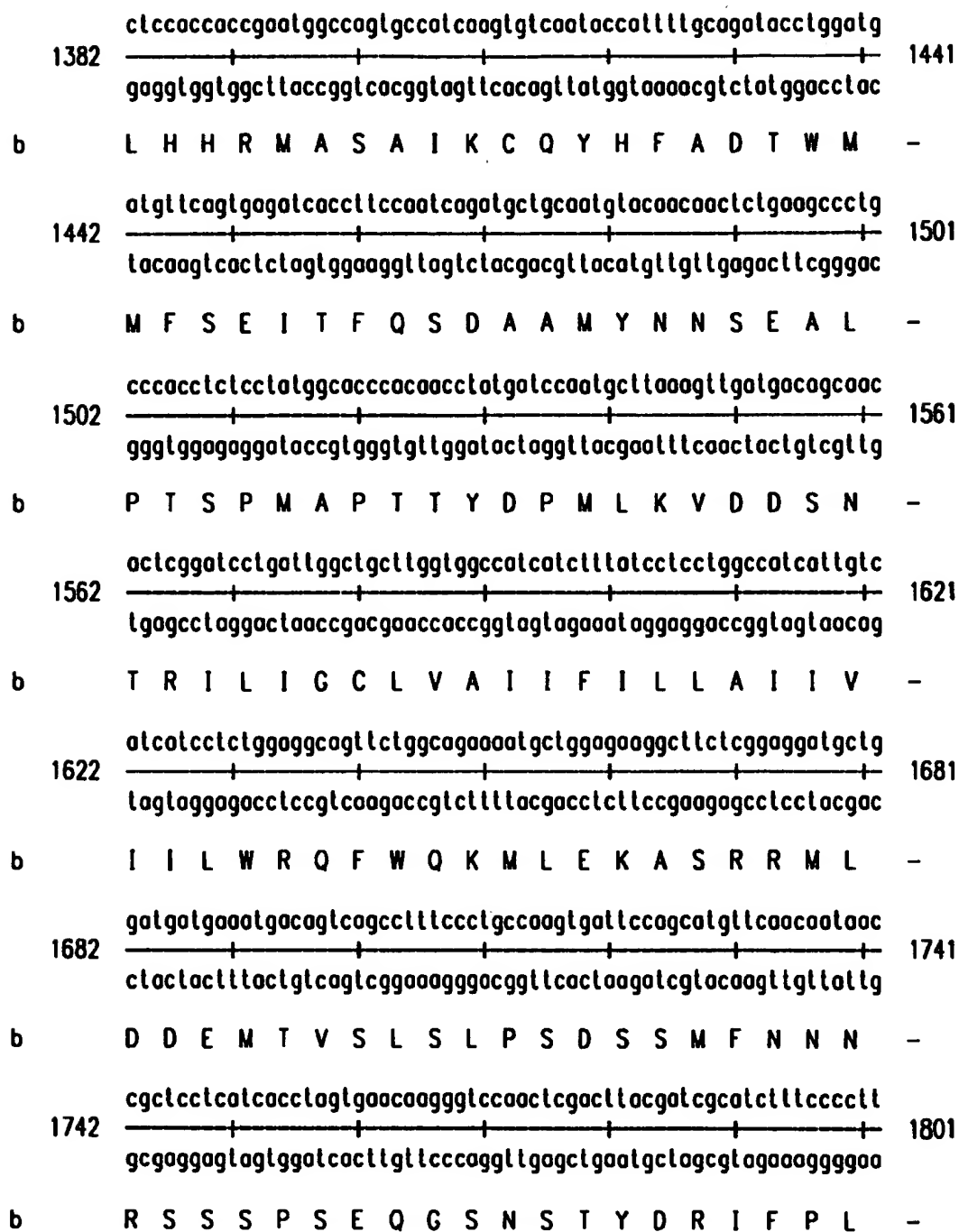
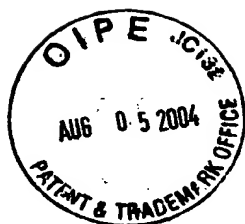


FIG.3D

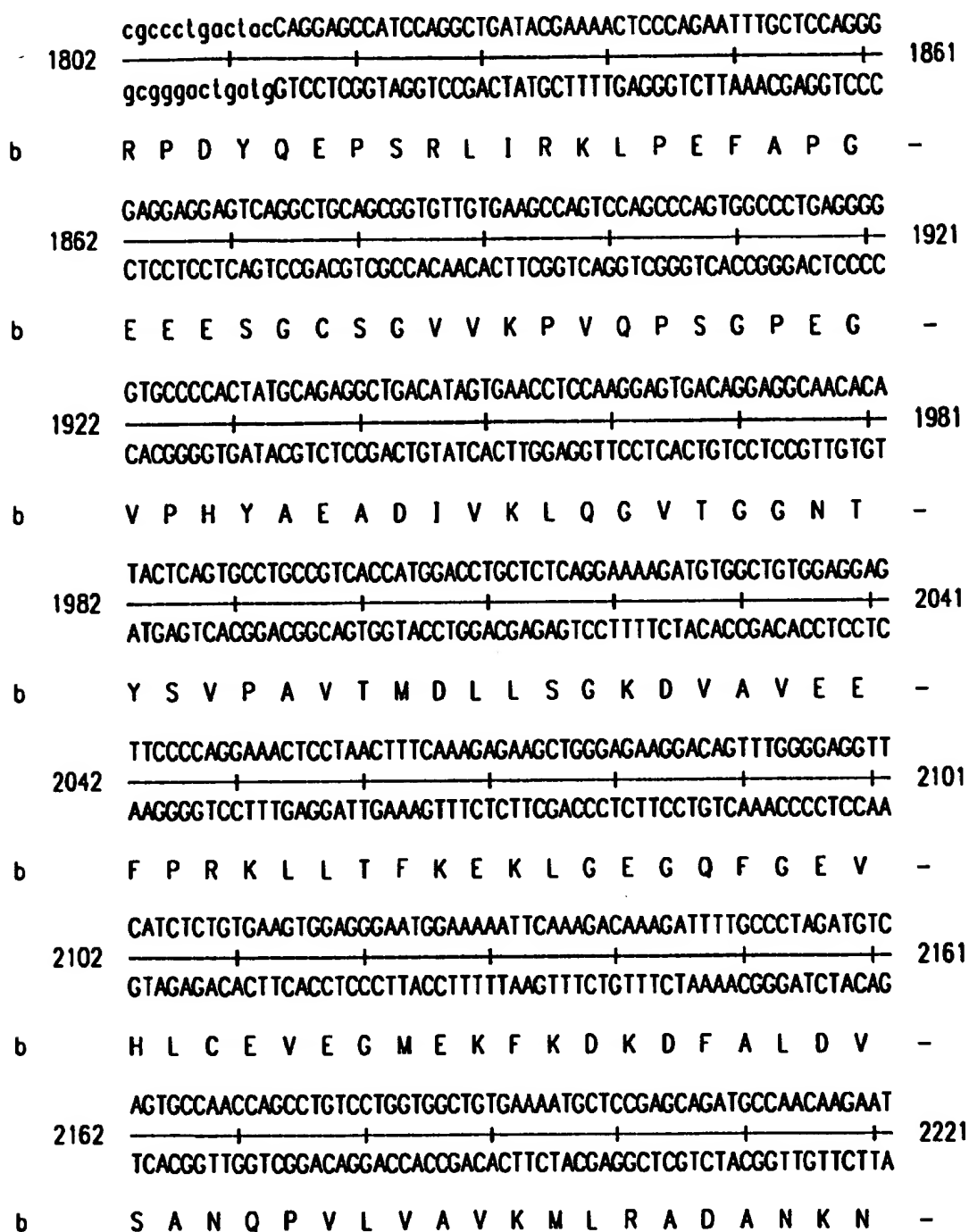


FIG.3E



2222 GCCAGGAATGATTTTCTTAAGGAGATAAAGATCATGTCTCGGCTCAAGGACCCAAACATC 2281
CGGTCCTTACTAAAAGAATTCTCTATTTCTAGTACAGAGCCGAGTTCTCGGGTTTGTAG

b A R N D F L K E I K I M S R L K D P N I -

2282 ATCCATCTATTAGCTGTGTGTATCACTGATGACCCTCTCTGTATGATCACTGAATACATG 2341
TAGGTAGATAATCGACACACATAGTACTACTGGGAGAGACATACTAGTGACTTATGTAC

b I H L L A V C I T D D P L C M I T E Y M -

2342 GAGAATGGAGATCTCAATCAGTTTCTTTCCGCCACGAGCCCCCTAATTCTTCTCCAGC 2401
CTCTTACCTCTAGAGTTAGTCAAAGAAAGGGCGGTCTCGGGGATTAAGAAGGAGGTGG

E N G D L N Q F L S R H E P P N S S S S -

2402 GATGTACGCACTGTCAGTTACCAATCTGAAGTTTATGGCTACCCAAATTGCCTCTGGC 2461
CTACATGCGTGACAGTCAATGTGGTTAGACTTCAAATACCGATGGGTTTAACGGAGACCG

b D V R T V S Y T N L K F M A T Q I A S G -

2462 ATGAAGTACCTTTCTCTCTTAATTTTGTTCACCGAGATCTGGCCACAGAACTGTTTA 2521
TACTTCATGGAAAGGAGAGAATTAACAAGTGGCTCTAGACCGGTGTGCTTTGACAAAT

b M K Y L S S L N F V H R D L A T R N C L -

2522 GTGGTAAGAACTACACAATCAAGATAGCTGACTTTGGAATGAGCAGGAACCTGTACAGT 2581
CACCCATTCTTGATGTGTAGTTCTATCGACTGAAACCTTACTCGTCCTTGGACATGTCA

b V G K N Y T I K I A D F G M S R N L Y S -

2582 GGTGACTATTACCGGATCCAGGGCCGGCAGTGCTCCCTATCCGCTGGATGTCTTGGGAG 2641
CCACTGATAATGGCCTAGGTCCCGGCCGTCACGAGGGATAGGCGACCTACAGAACCCTC

b G D Y Y R I Q G R A V L P I R W M S W E -

FIG.3F



FIG.3G



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b

3062 TTGCCCTCTTTTCTGGTCACCCCACTCCCTACCCCTGACTCATATATACTTTTTTTTT 3121
AACGGGAGAAAAGGACCAGTCGGGGTGAGGATGGGGACTGAGTATATGAAAAAAAAA

b

3122 TTACATTAAAGAACTAAAAAAAAAAAAAAAAAGGCG 3158
AATGTAATTTCTTGATTTTTTTTTTTTTTTTTTCCGC

b

FIG.3H



1	MILIPRMLLVLFLLLPILSSA...KAQVNPATCRYPLGMSGGQIPDEDIT	47 CCK-2
1	..MGPEALSSLLLLLVASGDADMKGHFDPAKCRYALGMQDRTIPSDIS	48 MCK-10
48	ASSQWSESTAAKYGRLDSEEGDGAWCPEIPVEPDDLKEFLQIDLHTLHF	97
49	ASSWSDSSTAARHSRLESSDGDGAWCAGSVFPKE.EEYLQVDLQRLHLV	97
98	TLVGTQGRRAGGCHGIEFAPMYKINYSRDGTRWISWRNRHGKQVLDGNSNP	147
98	ALVGTQGRHAGGLGKEFSRSYRLRYSRDGRRMGWKDRWGQEVISGNEP	147
148	YDIFLKDLEPPIVARFVRFIPVTDHSMNVCNRVELYGCWLDGLVSYNAP	197
148	EGVVLKDLGPPMVARLVRFYPRADRVMSVCLRVELYGCLWRDGLLSYAP	197
198	AGQQFVLPGGSIIYLNDVYDG.AVGYSMTEGLQLTQVSGLDQDFQTTH	246
198	VGQTMYLSEA..VYLNDSTYDGHTVGGLOYGGLQGLADGVVGLDDFRKSQ	245
247	EYHWPYDYGWRNESATNGYIEIMFEFDRIRNFTTMKVHCNMFAGKV	296
246	ELRVWPYDYGWSNHSFSSGYVEMEFEDRLRAFOAMQVHCNMMHTLGA	295
297	KIFKEVQC.YFRSEASEWVPNAISFPLVLDDVNP SARFVTVPLHHRMASA	345
296	RLPGGVECRFRRGPAWAGEPMRHNLCGNLGDPRARAVSVPLGGRVARF	345
346	IKQYHFADTMMFSEITFQSDAAMYNSEALPTS.....	380
346	LQCRFLFAGPWLLFSEISFISD.VVNNSSPALGGTFPPAPWMPGPPPTN	394
381PMAPTYYDPMKVDSDNTRILIGCLVATIFILLATIVITLWRQFWQ	426 TRANSMEMBRANE
395	FSSLELEPRGQOPVAKAEGSPTAILIGCLVATILLLLTIALMLWRLHWR	444 REGION
427	KMLEKASRRMLDDEMTVSLSLPSDSSMFMNNRSSSPSEQGSNSTYDRIFP	476
445	RLLSKAERRVLEEELTVHLSVPGDTILINRPGPREP.....	481

FIG.4A

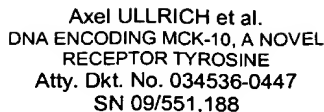
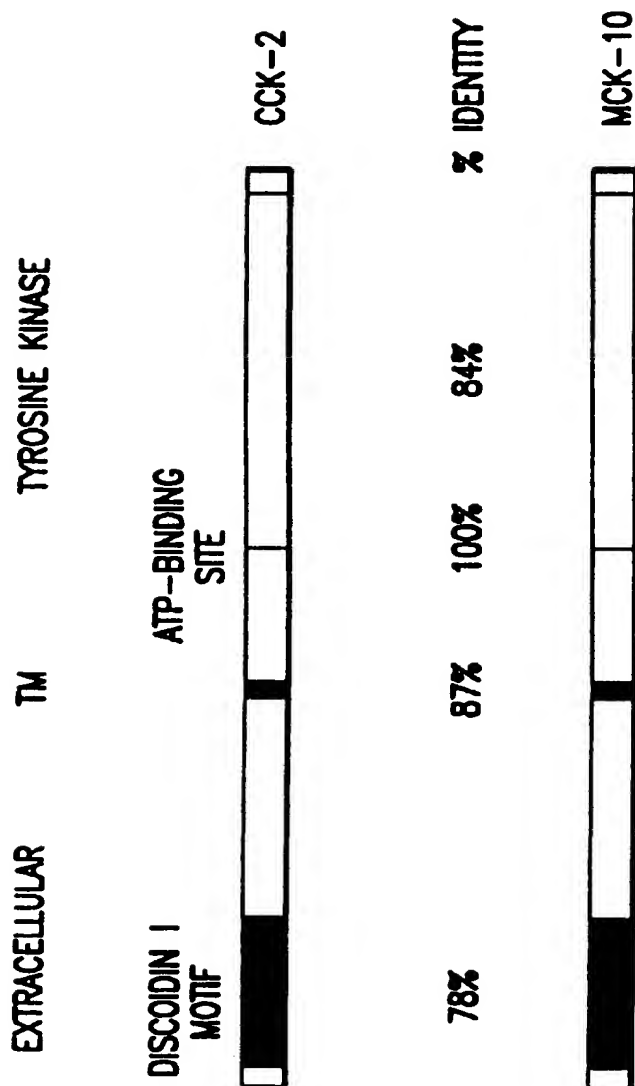
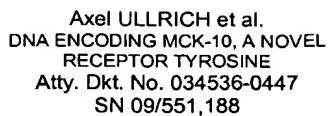


FIG. 4B





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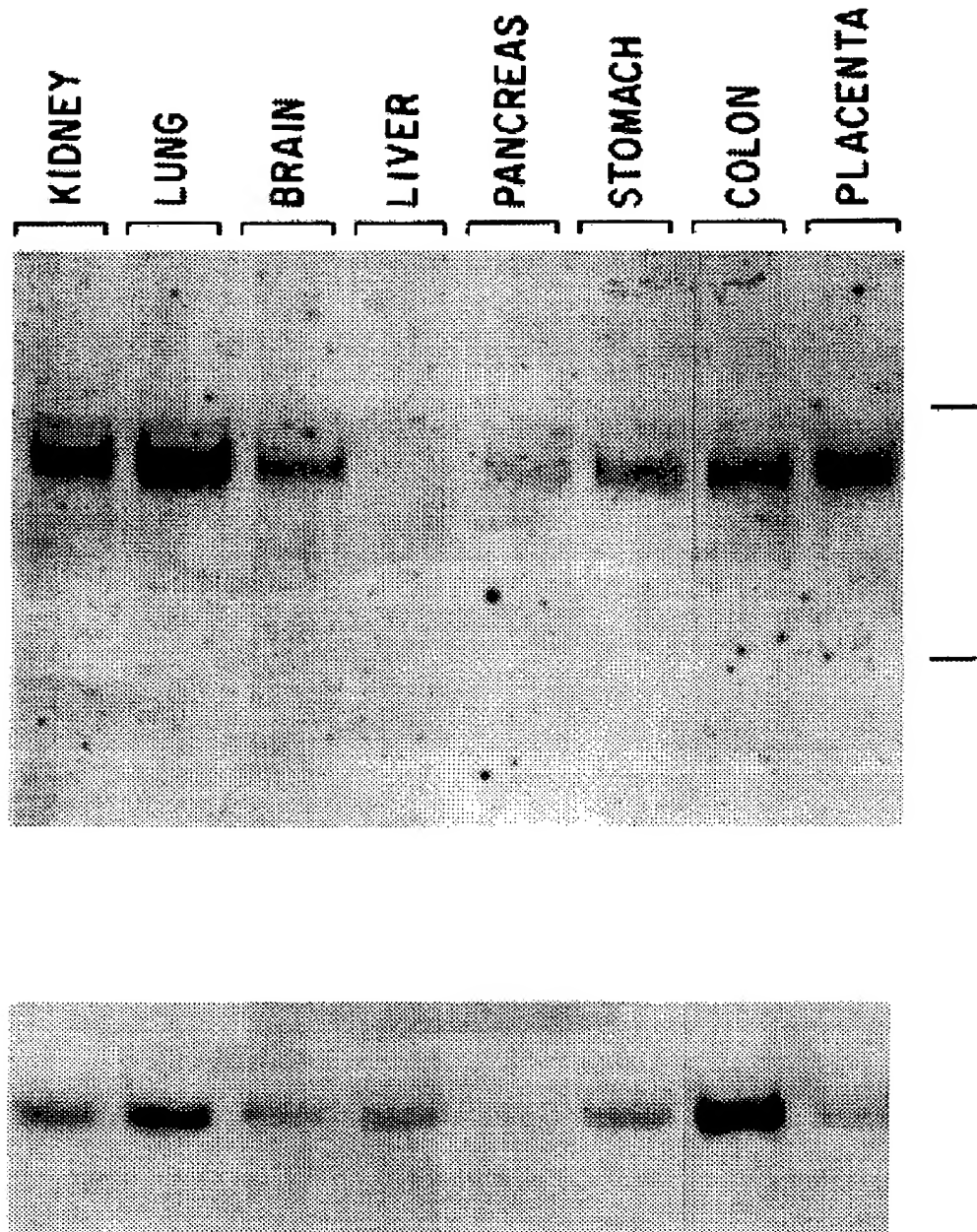


FIG.5A

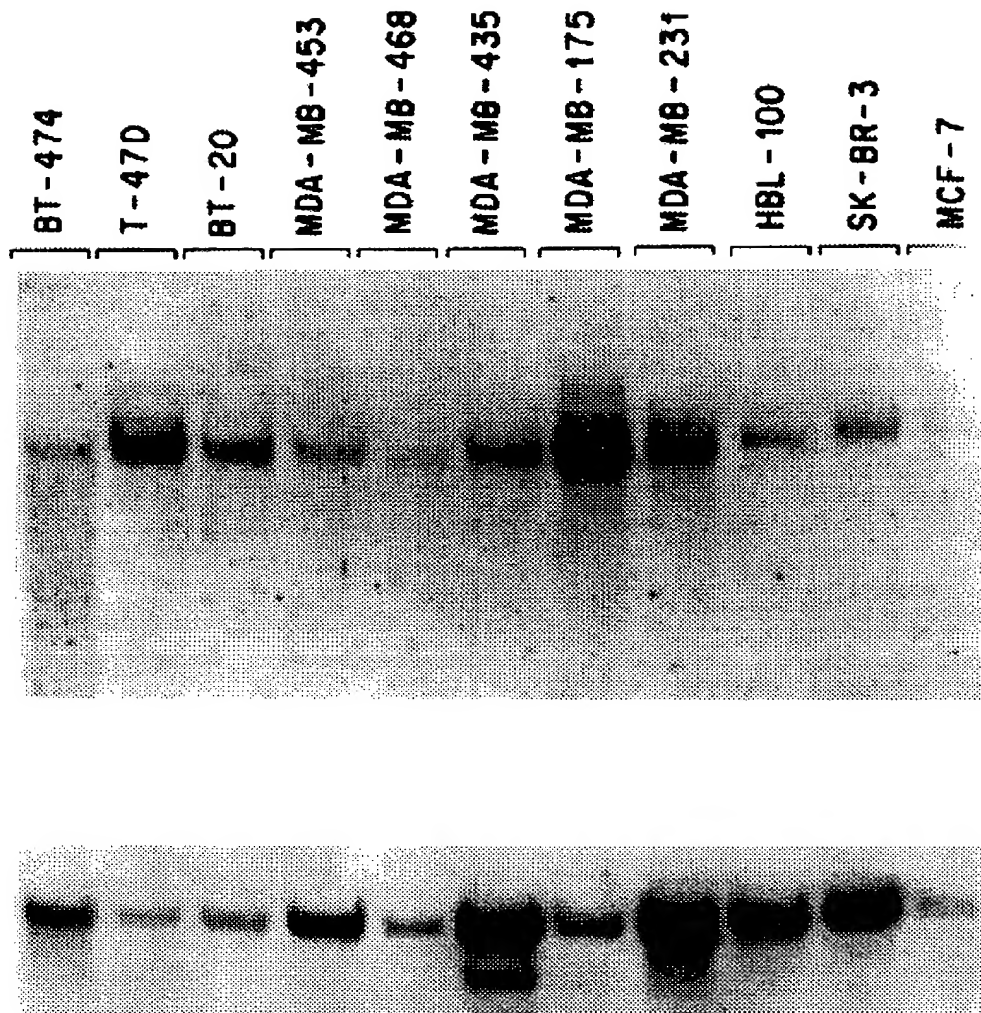


FIG.5B

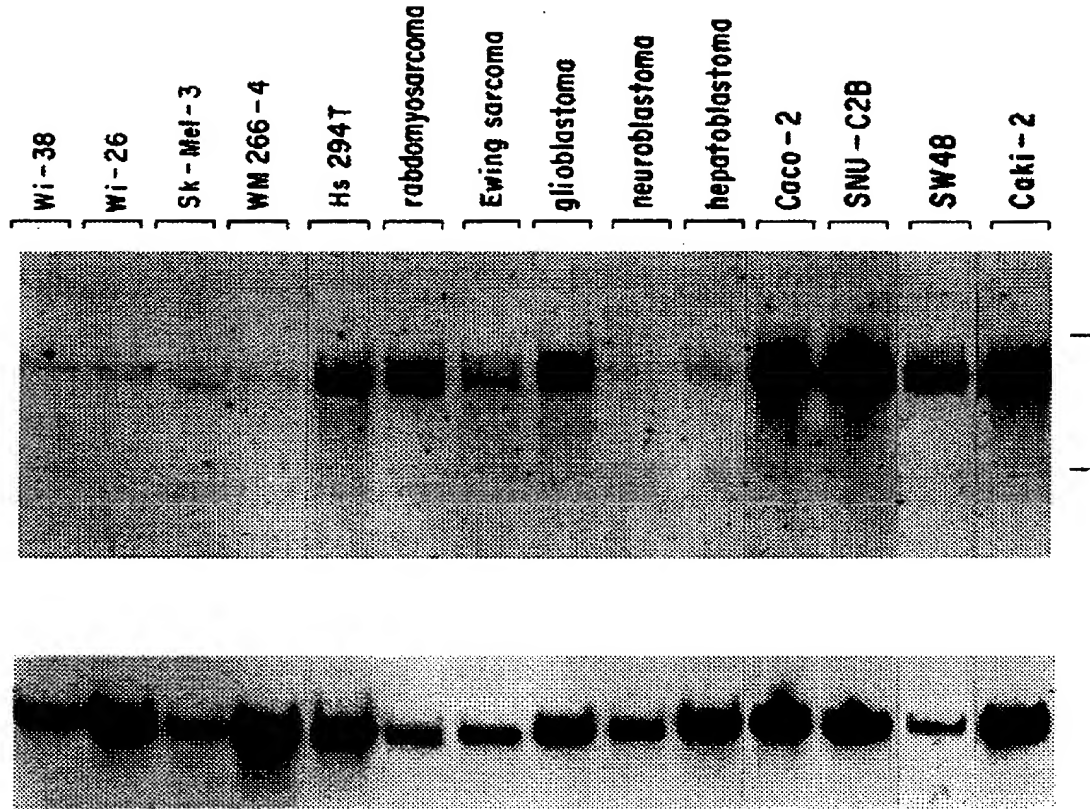


FIG.5C

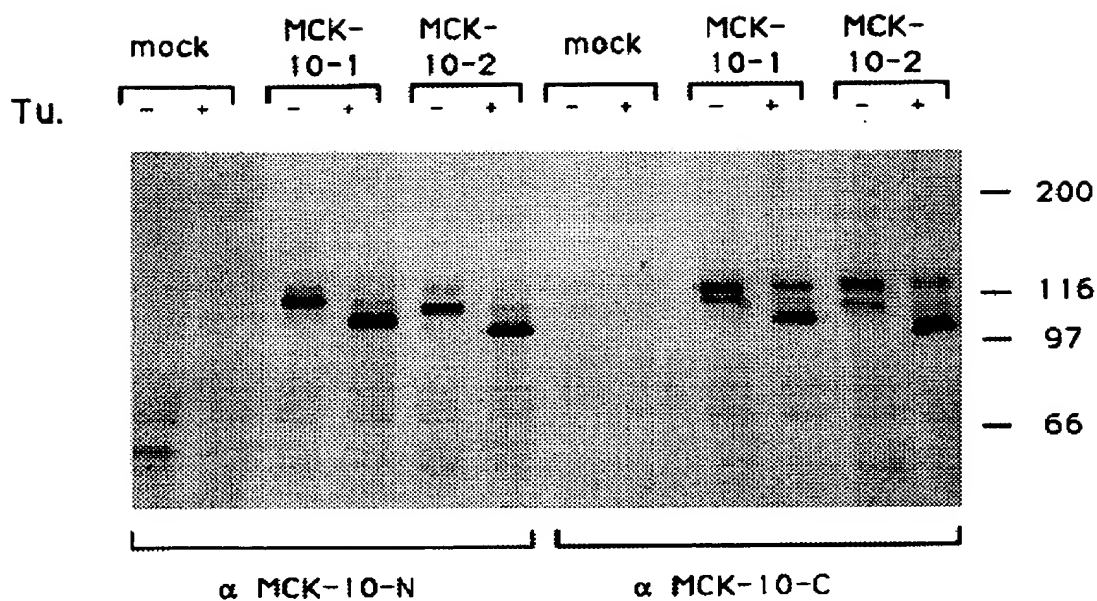
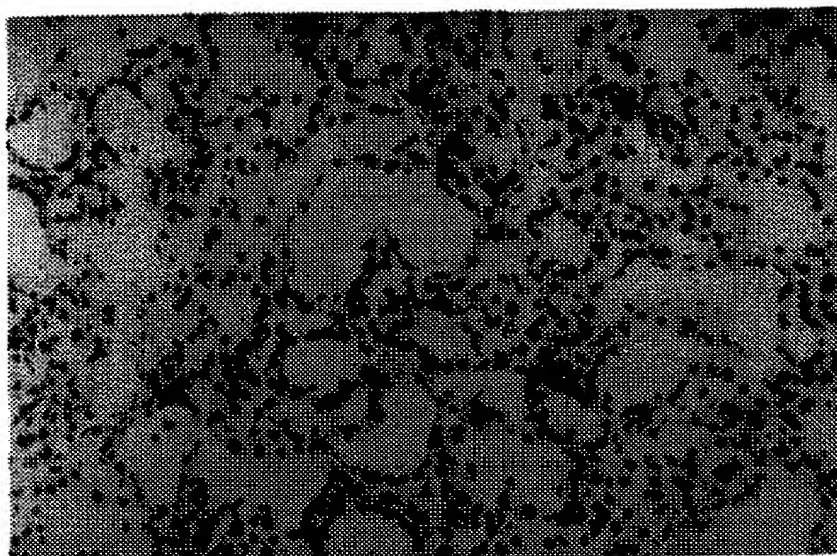


FIG.6B

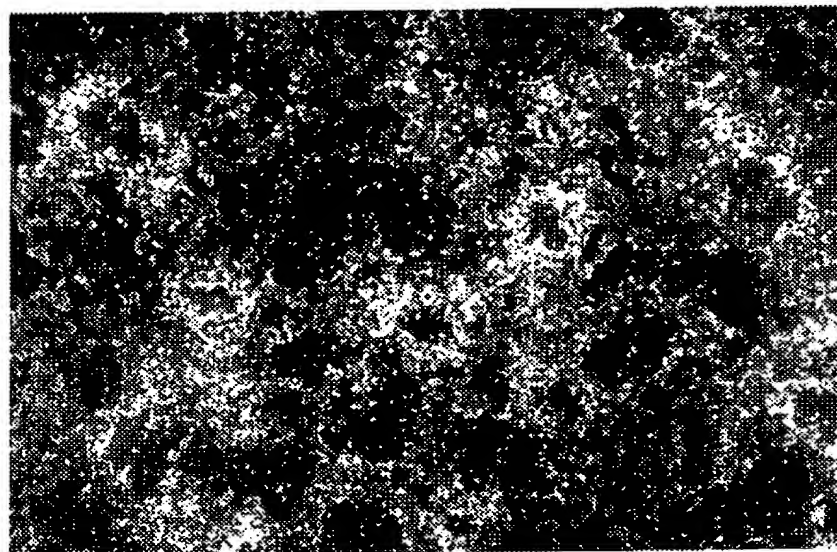


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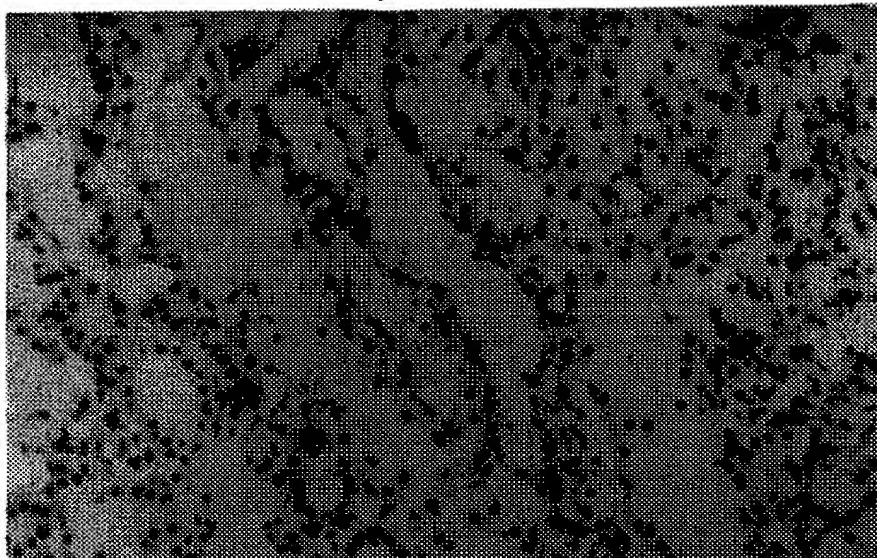
Lightfield

FIG.7A



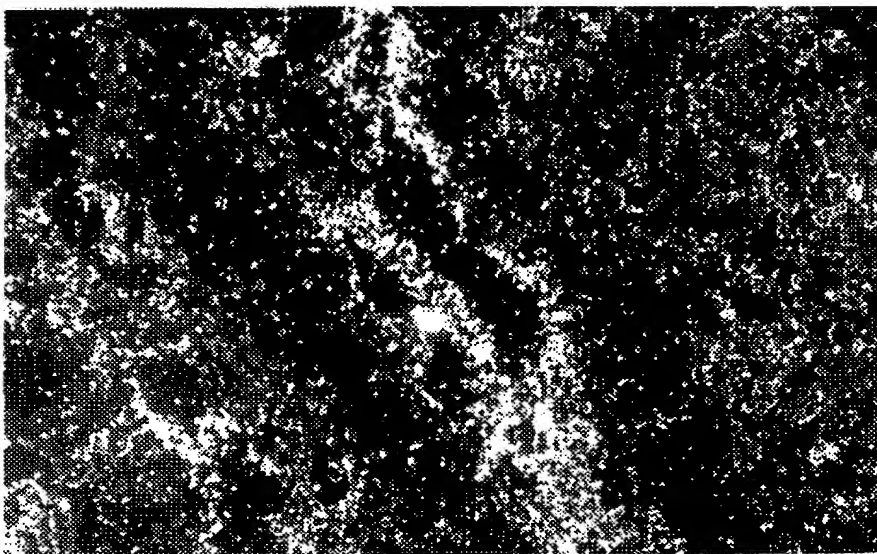
Darkfield

FIG.7B



Lightfield

FIG.8A

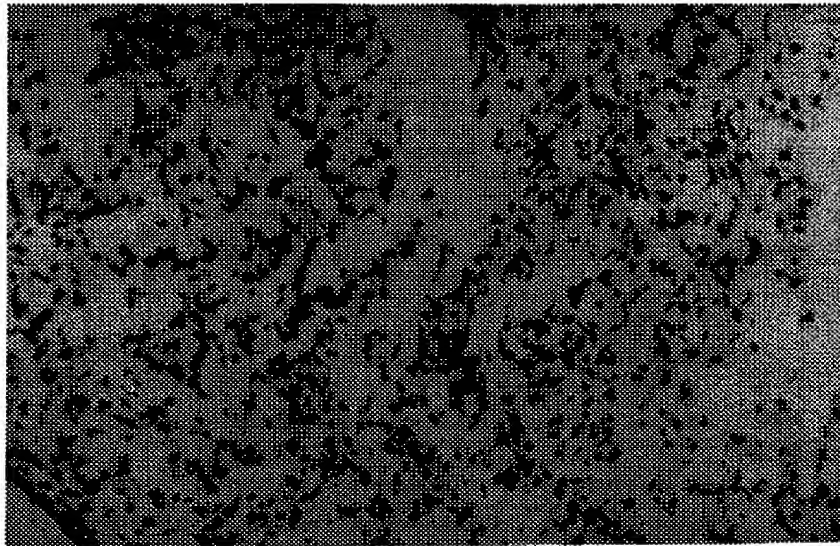


Darkfield

FIG.8B

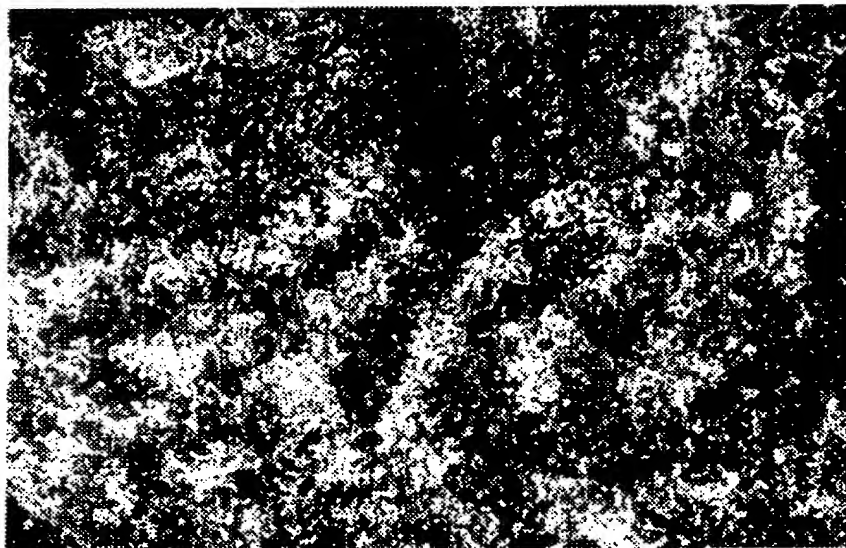


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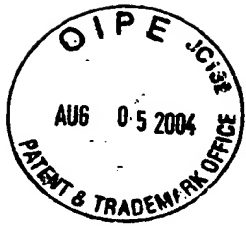
Lightfield

FIG.9A

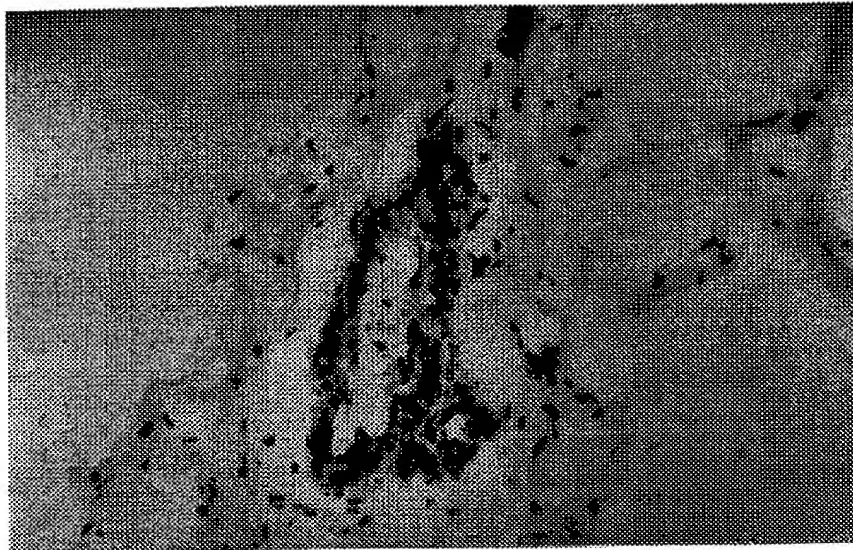


Darkfield

FIG.9B

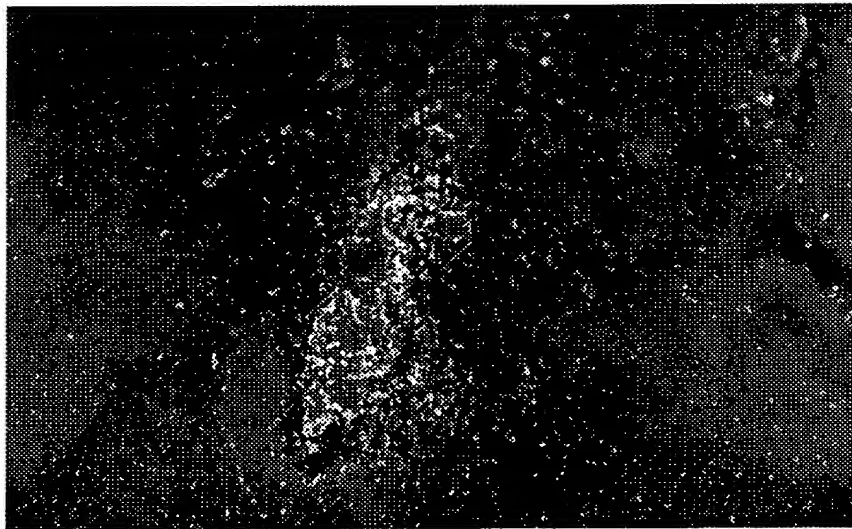


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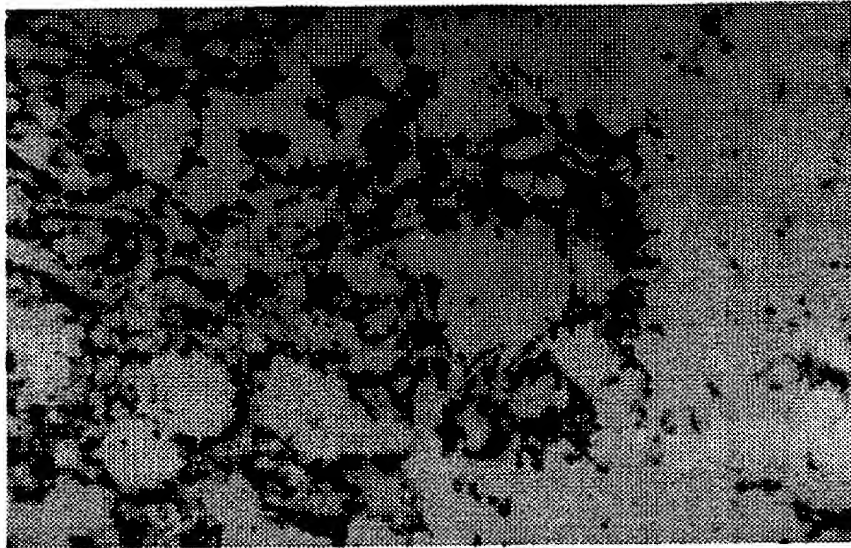
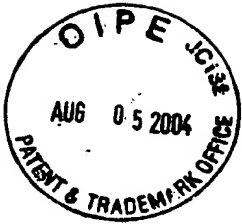
Lightfield

FIG.10A



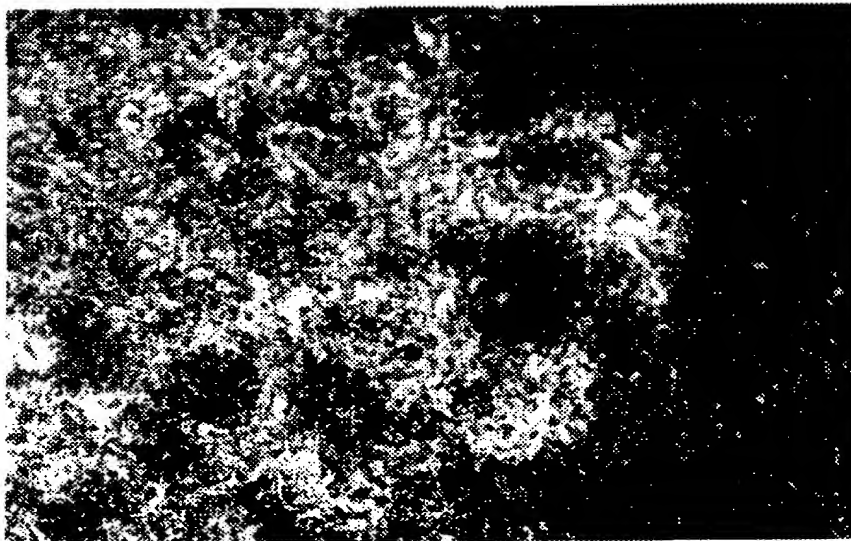
Darkfield

FIG.10B



Lightfield

FIG.11A

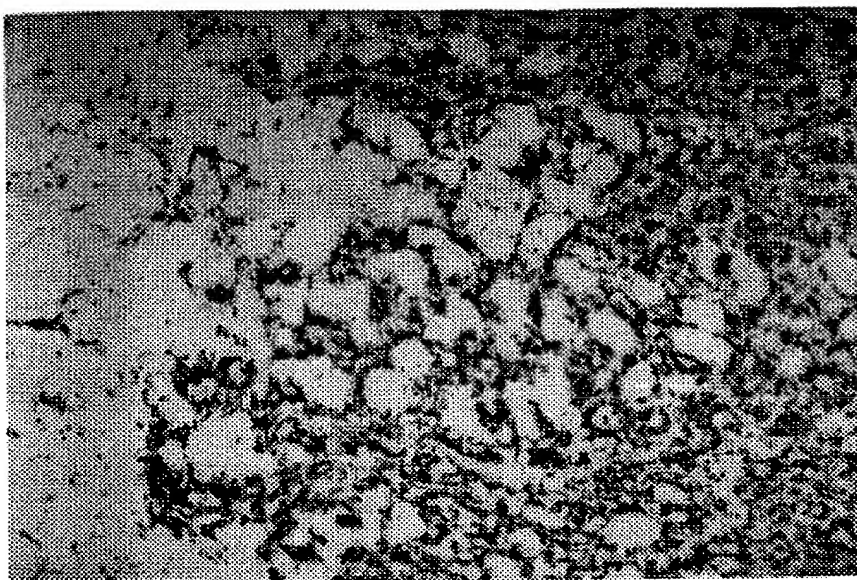


Darkfield

FIG.11B

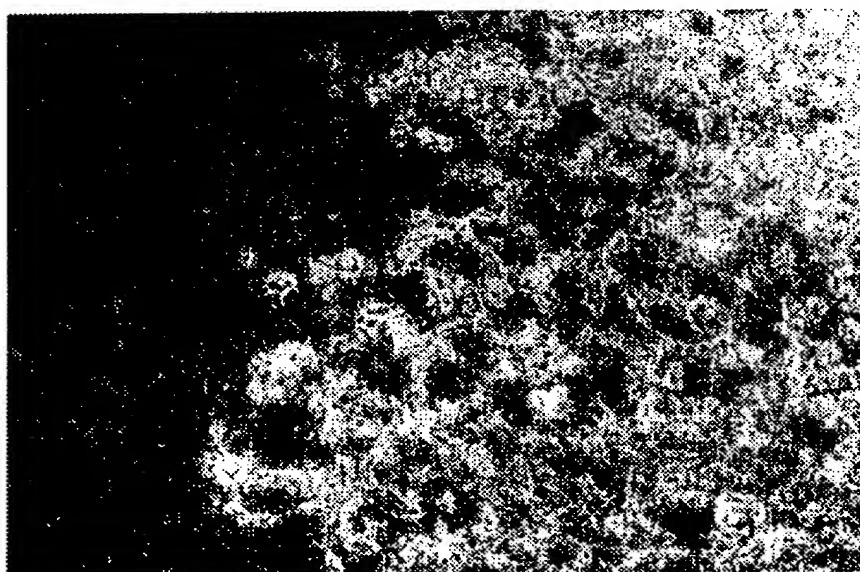


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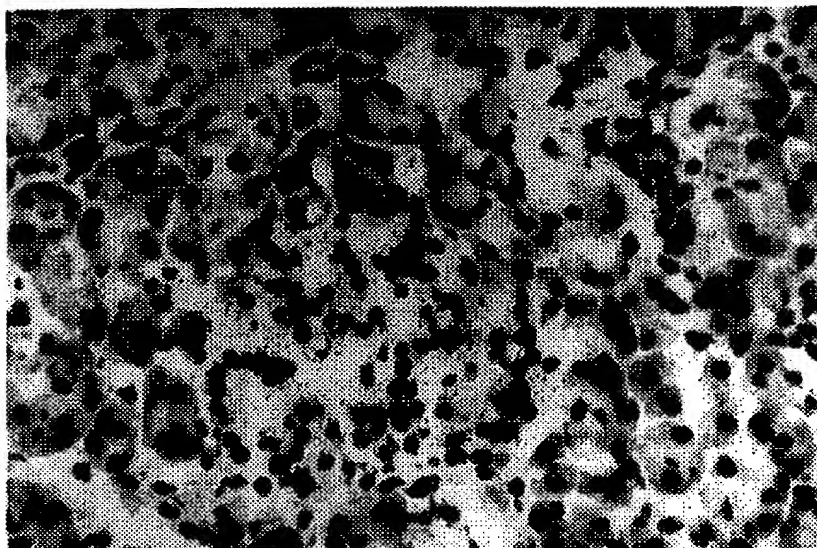
Lightfield

FIG.12A



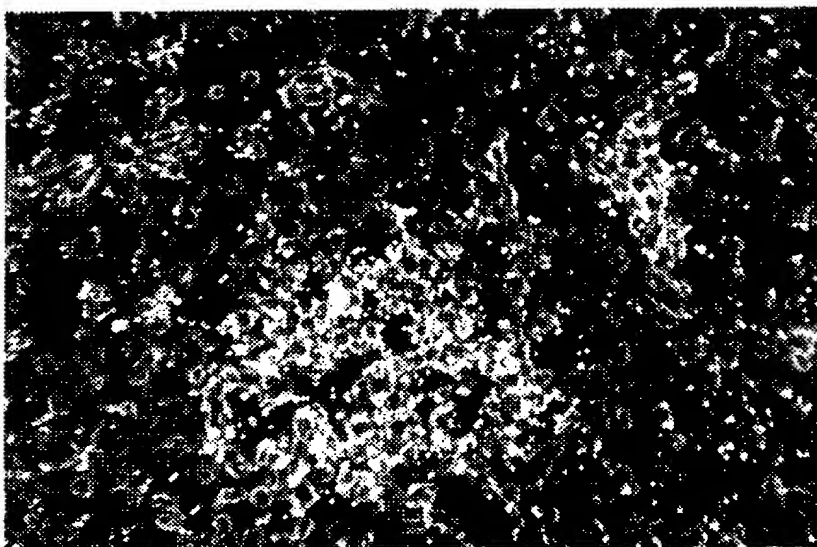
Darkfield

FIG.12B



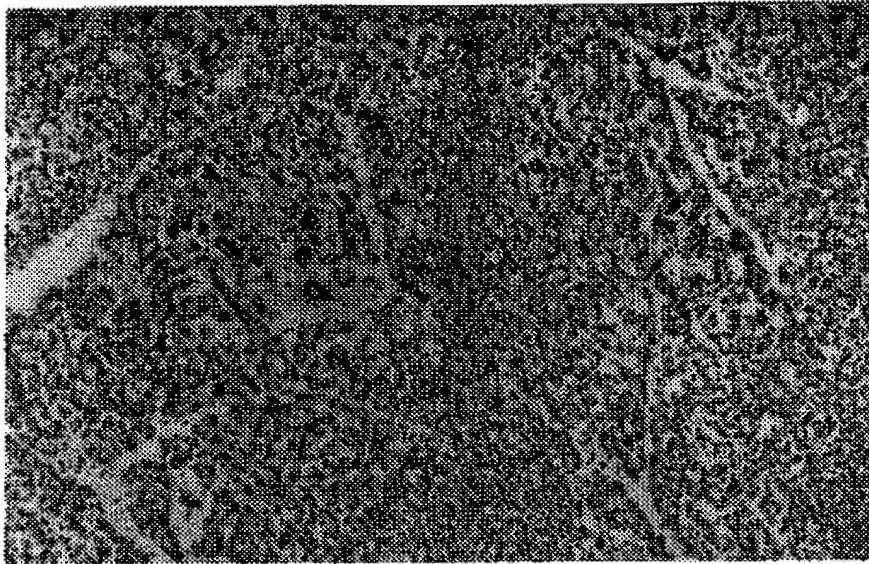
Lightfield

FIG.13A



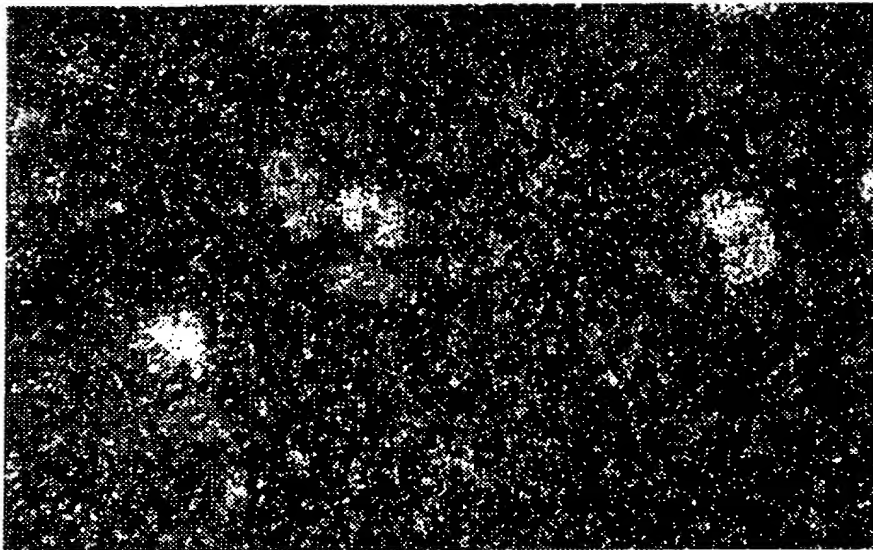
Darkfield

FIG.13B



Lightfield

FIG. 14A



Darkfield

FIG. 14B

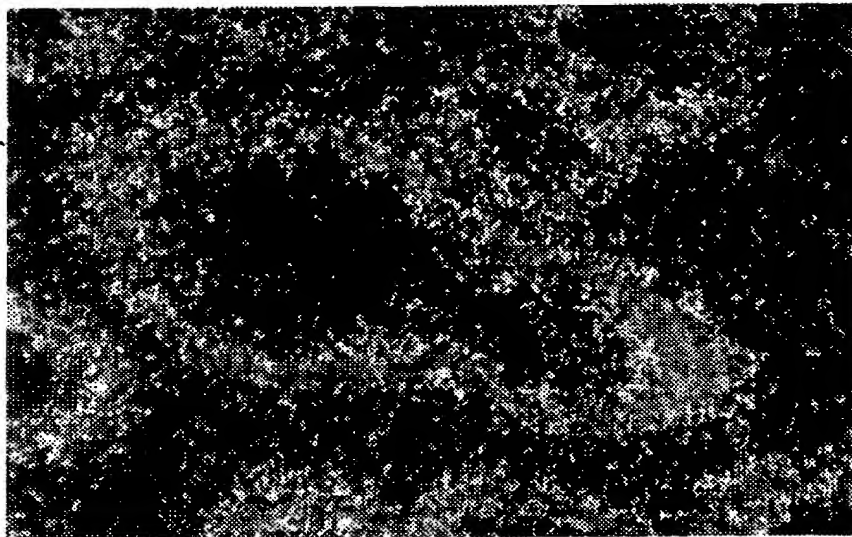


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Lightfield

FIG.15A

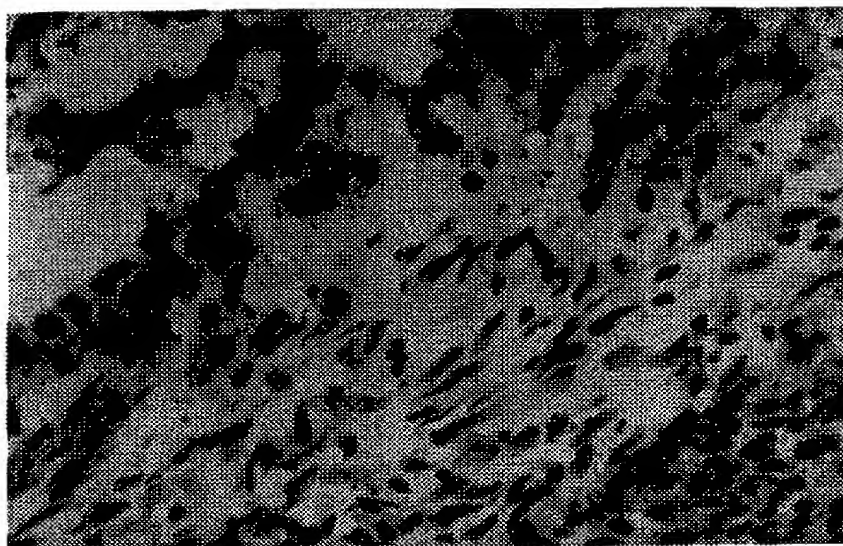


Darkfield

FIG.15B

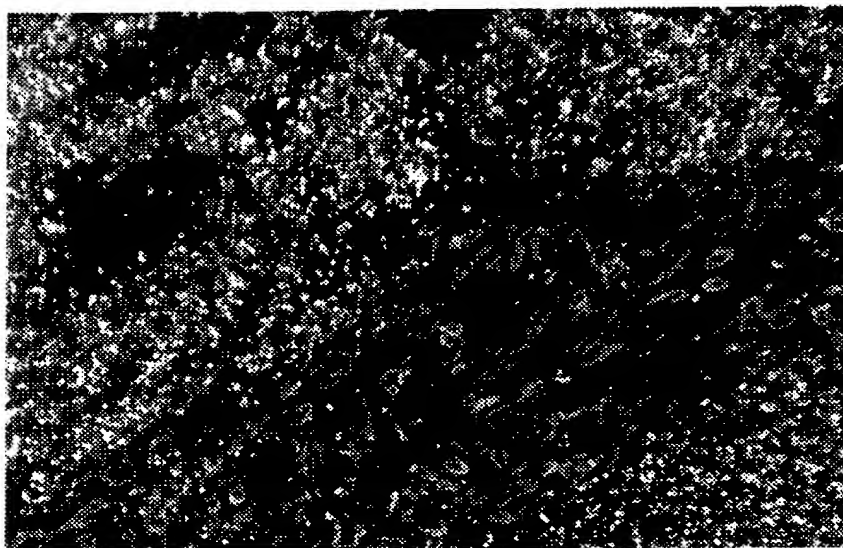


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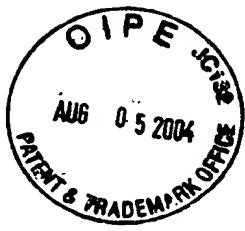
Lightfield

FIG.16A



Darkfield

FIG.16B

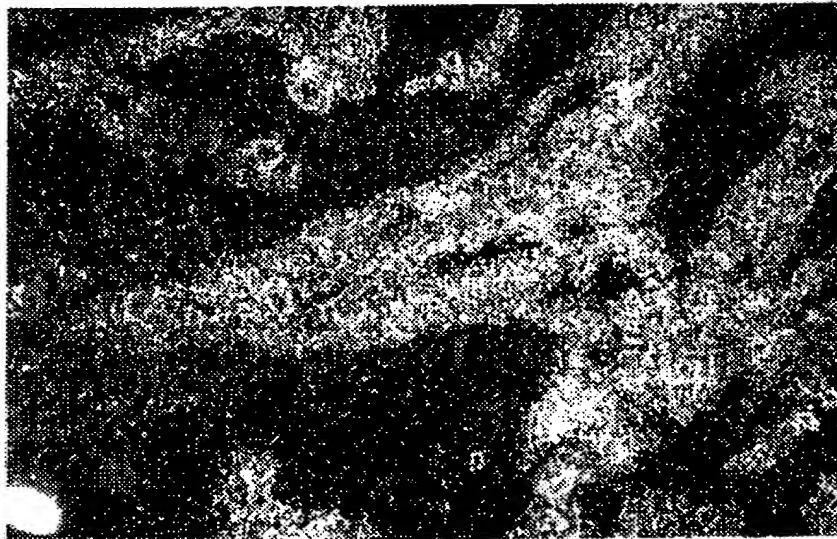


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Lightfield

FIG.17A



Darkfield

FIG.17B

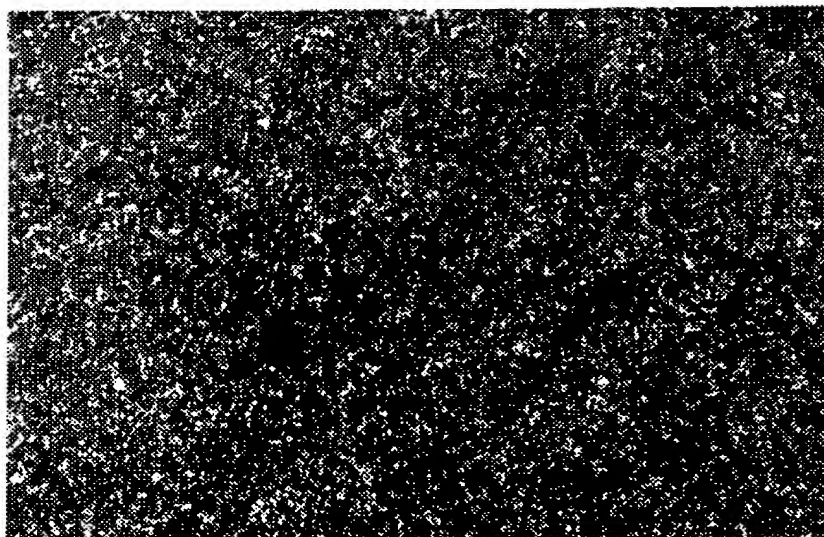


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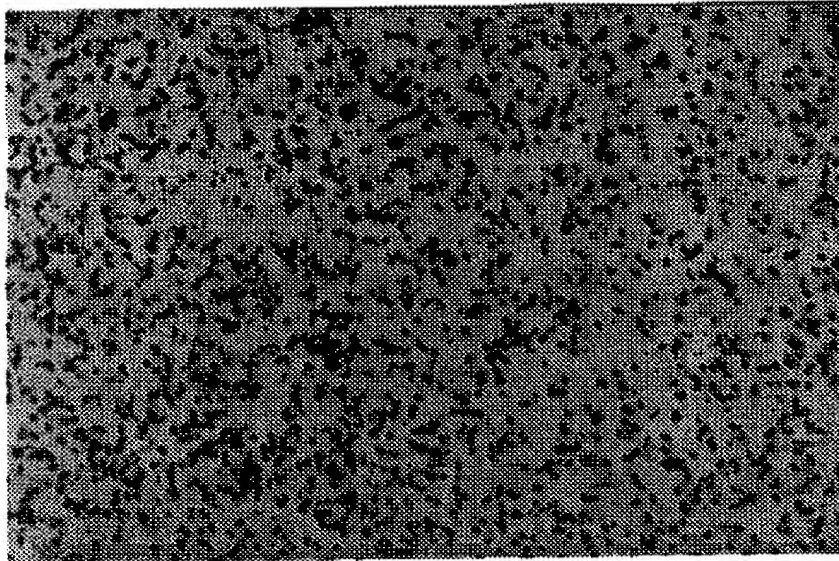
Lightfield

FIG.18A



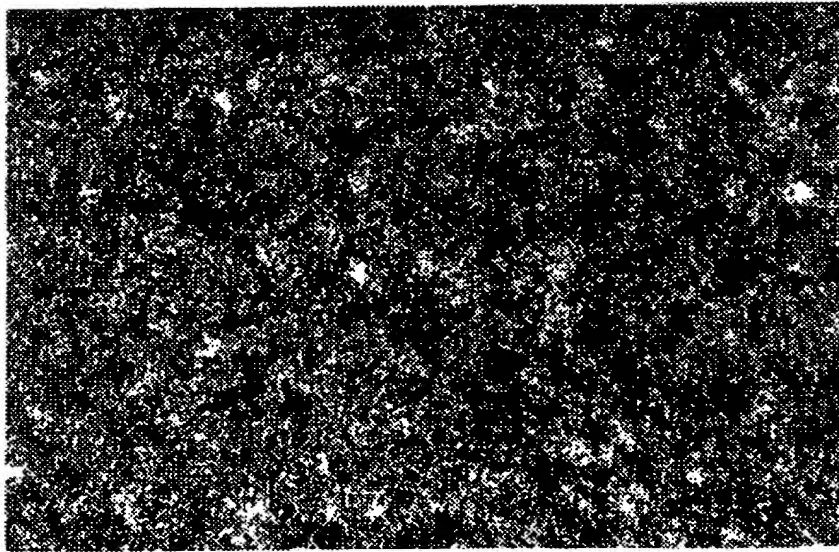
Darkfield

FIG.18B



Lightfield

FIG.19A

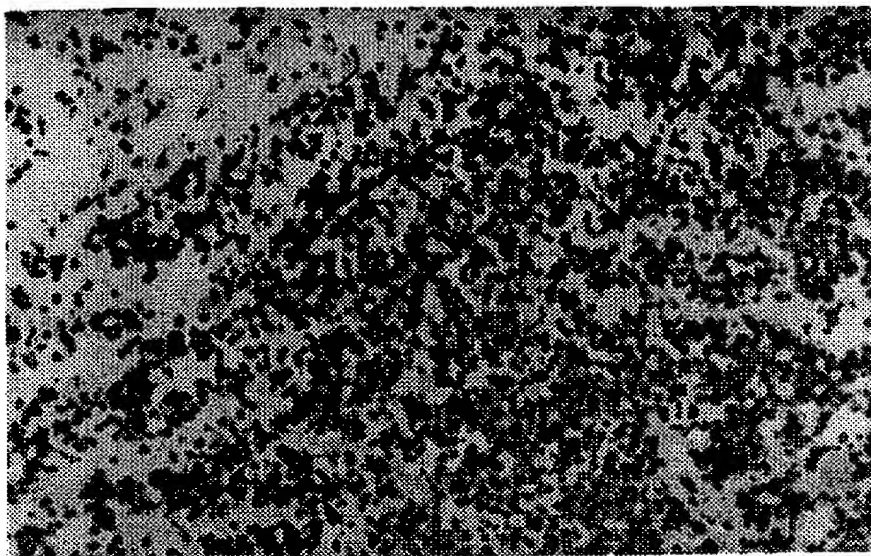


Darkfield

FIG.19B

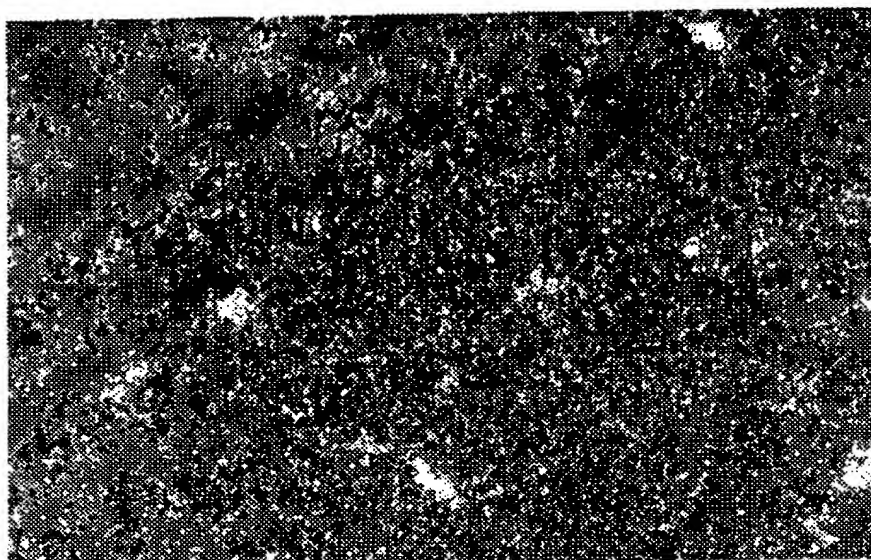


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Lightfield

FIG.20A

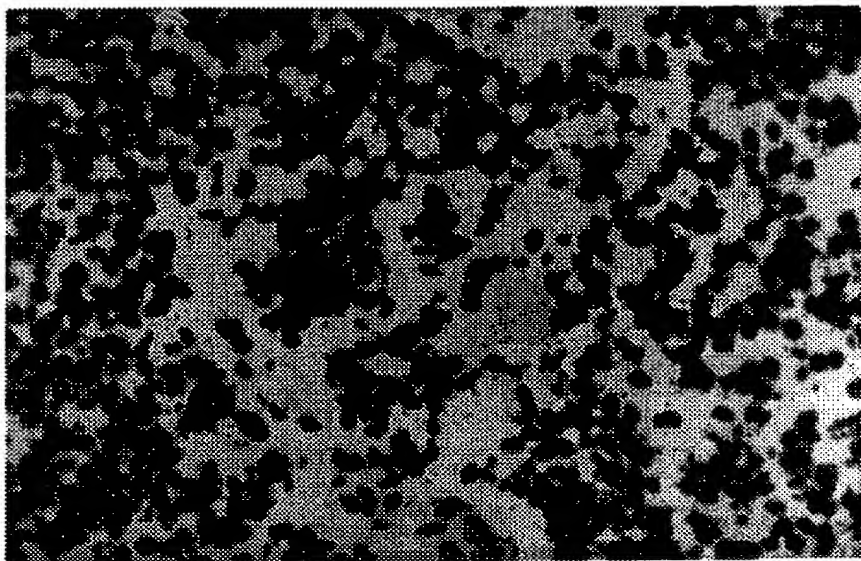


Darkfield

FIG.20B

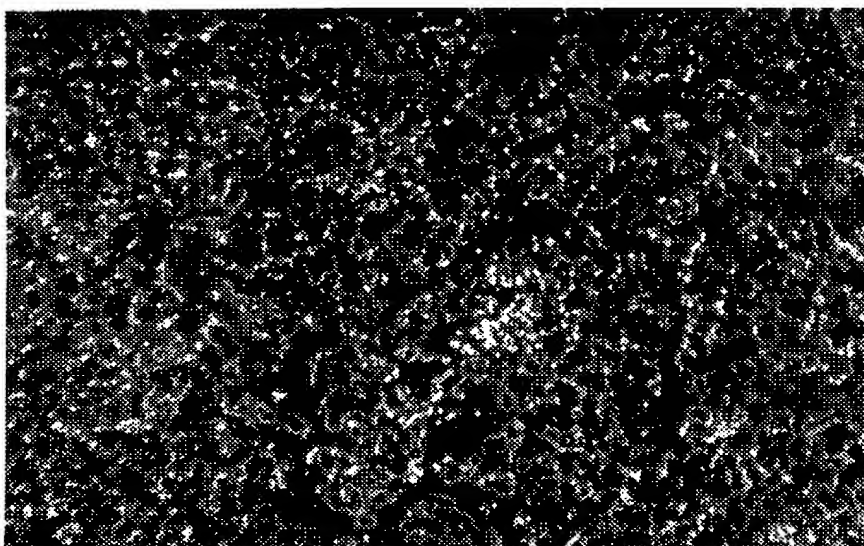


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Lightfield

FIG.21A



Darkfield

FIG.21B